

GenCore version 5.1.6
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OM protein - protein search, using sw model

Run on: July 1, 2003, 08:41:43 ; Search time 15 Seconds
 (without alignments)
 435,459 Million cell updates/sec

Title: US-09-508-710-2

Perfect score: 1173

Sequence: 1 MAGEKGIVLDFWVSPFGQR.....YSPDKVYDFIGLLKKYKIE 222

Scoring table: BLOSUM62

Gapop 10.0 , Gapext 0.5

Searched: 262574 seqs, 29422922 residues

Total number of hits satisfying chosen parameters: 262574

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
 Maximum Match 100%
 Listing first 45 summaries

Database : Issued_Patents_AA:
 1: /cgn2_6/ptodata/1/iaa/5A_COMB.pep:
 2: /cgn2_6/ptodata/1/iaa/5B_COMB.pep:
 3: /cgn2_6/ptodata/1/iaa/6A_COMB.pep:
 4: /cgn2_6/ptodata/1/iaa/6B_COMB.pep:
 5: /cgn2_6/ptodata/1/iaa/PCTUS_COMBO.pep:
 6: /cgn2_6/ptodata/1/iaa/backfiles1.pep:
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 *
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Prod. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query	Match	Length	DB	ID	Description
1	857	73.1	224	2	US-09-247-759-22		Sequence 22, Appl
2	857	73.1	224	3	US-09-248-335-22		Sequence 22, Appl
3	818.5	69.8	225	2	US-08-924-759-12		Sequence 12, Appl
4	818.5	69.8	225	3	US-09-248-335-12		Sequence 12, Appl
5	683.5	58.3	219	3	US-08-924-747-6		Sequence 6, Appl
6	683.5	58.3	219	4	US-09-296-715-6		Sequence 6, Appl
7	681	58.1	218	4	US-09-247-738-6		Sequence 6, Appl
8	664.5	56.6	219	3	US-08-924-747-20		Sequence 20, Appl
9	664.5	56.6	219	4	US-09-247-738-20		Sequence 20, Appl
10	664.5	56.6	219	4	US-09-296-715-20		Sequence 20, Appl
11	639	54.5	216	3	US-08-924-747-1		Sequence 10, Appl
12	639	54.5	216	4	US-09-247-738-12		Sequence 12, Appl
13	639	54.5	216	4	US-09-296-715-12		Sequence 12, Appl
14	601.5	51.3	200	3	US-08-924-747-18		Sequence 18, Appl
15	601.5	51.3	200	4	US-09-247-738-20		Sequence 18, Appl
16	601.5	51.3	200	4	US-09-296-715-18		Sequence 18, Appl
17	566.5	48.3	221	4	US-09-247-738-44		Sequence 44, Appl
18	488	41.6	230	3	US-09-248-335-44		Sequence 44, Appl
19	480	40.9	147	1	US-08-525-507-2		Sequence 2, Appl
20	432.5	36.9	237	3	US-09-248-335-48		Sequence 48, Appl
21	431.5	36.8	225	3	US-09-248-335-40		Sequence 40, Appl
22	430	36.7	228	3	US-09-248-335-70		Sequence 70, Appl
23	426.5	36.4	231	3	US-09-248-335-54		Sequence 54, Appl
24	406	34.6	236	3	US-09-248-335-74		Sequence 74, Appl
25	401	34.2	238	3	US-09-248-335-38		Sequence 38, Appl
26	399.5	34.1	233	3	US-09-248-335-60		Sequence 60, Appl
27	389.5	33.2	235	3	US-08-924-747-24		Sequence 24, Appl

ALIGNMENTS

RESULT 1	US-08-924-759-22	Sequence 22, Application US/08924759
	; Patent No. 596229	GENERAL INFORMATION:
	; APPLICANT: MCGONIGLE, BRIAN	APPLICANT: O'KEEFE, DANIEL
	; TITLE OF INVENTION: PLANT GLUTATHIONE-S-TRANSFERASE	TITLE OF INVENTION: ENYMES
	; NUMBER OF SEQUENCES: 24	CORRESPONDENCE ADDRESS:
	; ADDRESSEE: E. I. DU PONT DE NEMOURS AND COMPANY	CITY: 1007 MARKET STREET
	; STATE: DELAWARE	STATE: WILMINGTON
	; COUNTRY: UNITED STATES OF AMERICA	COUNTRY: UNITED STATES OF AMERICA
	; ZIP: 19898	ZIP: 19898
	COMPUTER READABLE FORM:	COMPUTER READABLE FORM:
	MEDIUM TYPE: DISKETTE, 3.5 INCH	MEDIUM TYPE: DISKETTE, 3.5 INCH
	COMPUTER: IBM PC COMPATIBLE	COMPUTER: IBM PC COMPATIBLE
	OPERATING SYSTEM: MICROSOFT WORD FOR WINDOWS 95	OPERATING SYSTEM: MICROSOFT WORD FOR WINDOWS 95
	SOFTWARE: MICROSOFT WORD VERSION 7.0A	SOFTWARE: MICROSOFT WORD VERSION 7.0A
	CURRENT APPLICATION DATA:	CURRENT APPLICATION DATA:
	APPLICATION NUMBER: US/08/924,759	APPLICATION NUMBER: US/08/924,759
	FILING DATE:	FILING DATE:
	CLASSIFICATION: 435	CLASSIFICATION: 435
	ATTORNEY/AGENT INFORMATION:	ATTORNEY/AGENT INFORMATION:
	NAME: FLOYD, LINDA AXAMTHY	NAME: FLOYD, LINDA AXAMTHY
	REFERENCE/DOCKET NUMBER: CL-1128	REFERENCE/DOCKET NUMBER: CL-1128
	TELECOMMUNICATION INFORMATION:	TELECOMMUNICATION INFORMATION:
	TELEPHONE: 302-892-8112	TELEPHONE: 302-892-8112
	TELEFAX: 302-771-0164	TELEFAX: 302-771-0164
	INFO FOR SEQ ID NO: 22:	INFO FOR SEQ ID NO: 22:
	SEQUENCE CHARACTERISTICS:	SEQUENCE CHARACTERISTICS:
	LENGTH: 224 amino acids	LENGTH: 224 amino acids
	TYPE: amino acid	TYPE: amino acid
	STRANDEDNESS: not relevant	STRANDEDNESS: not relevant
	TOPOLOGY: not relevant	TOPOLOGY: not relevant
	MOLECULE TYPE: protein	MOLECULE TYPE: protein
	ORIGINAL SOURCE:	ORIGINAL SOURCE:
	SEQUENCE TYPE: MAIZE	SEQUENCE TYPE: MAIZE
	IMMEDIATE SOURCE:	IMMEDIATE SOURCE:
	CLONE: CEB5.PR0049.A11	CLONE: CEB5.PR0049.A11
US-08-924-759-22	Query Match	73.1%; Score 857; DB 2; Length 224;
	Best Local Similarity	72.9%; Prod. No. 9.7e-91;
	Matches 164; Conservative 27; Mismatches 30; Indels 4; Gaps 3;	Matches 164; Conservative 27; Mismatches 30; Indels 4; Gaps 3;

QY 1 MAGEK - GLVLLDFWVSPFGQRVRLAEGKGLPVEYAEEDLMAGSKDRLLRANPVHKIP 58
 1 ||| 1 ||| 1 ||| 1 ||| 1 ||| 1 ||| 1 ||| 1 ||| 1 : 1 ||| 1 ||| 1 ||| 1 ||| 1 |||
 1 MAEKKQGLQQLDFWVSPFGQRCLRLDEKGLAYELEQDL - RNSSELLRANPVHKIP 59
 1 ||| 1 ||| 1 ||| 1 ||| 1 ||| 1 ||| 1 ||| 1 ||| 1 ||| 1 : 1 ||| 1 ||| 1 ||| 1 ||| 1 |||
 59 VLLHDGRAVNESLIILQYLEAEPD - APALLPSDPTYARAQARFWADYVDKKYDCCGSLRW 117
 1 ||| 1 ||| 1 ||| 1 ||| 1 ||| 1 ||| 1 ||| 1 ||| 1 ||| 1 ||| 1 : 1 ||| 1 ||| 1 ||| 1 ||| 1 |||
 Db 60 VLLHDGRPVCLSLVQYLDEAEPAPALLPADPPTYARAQARFWADYVDKLYDCGTRLW 119
 118 KLKGEPQAQAAEMLDILKTDGALDKPFGGDKGTFVDAAFAPPTAWPHSYERGEFS 177
 1 ||| 1 ||| 1 ||| 1 ||| 1 ||| 1 ||| 1 ||| 1 : 1 ||| 1 ||| 1 ||| 1 ||| 1 ||| 1 ||| 1 |||
 Db 120 KLKGDSQAQAAEMVYLRLTLEGALGDPFGGDKGTFVDAALVPTSWFLAYDRGGVS 179
 1 ||| 1 ||| 1 ||| 1 ||| 1 ||| 1 ||| 1 ||| 1 : 1 ||| 1 ||| 1 ||| 1 ||| 1 ||| 1 ||| 1 |||

QY 178 LPEVAKIAAWAKRCGEREVSAKSLSPDKYDFICLLKKYGLIE 222
 1 :
 Db 180 VEKECPRLAAWAKRCERPSVAKNLYPPEKTYDFVGMKRLGIE 224

RESULT 2

US-09-248-335-22

; Sequence 22, Application US/09248335

; GENERAL INFORMATION:

; APPLICANT: MCCONIGLE, BRIAN

; TITLE OF INVENTION: PLANT GLUTATHIONE-S-TRANSFERASE ENZYMES

; FILE REFERENCE: CL-1128-A

; CURRENT APPLICATION NUMBER: US/09/248,335

; CURRENT FILING DATE: 1999-02-10

; EARLIER APPLICATION NUMBER: 08/924,759

; EARLIER FILING DATE: 1997-September-05

; NUMBER OF SEQ ID NOS: 74

; SOFTWARE: Microsoft Word Version 7.0A

; SEQ ID NO: 22

; TYPE: PRT

; ORGANISM: maize

US-09-248-335-22

Query Match 73.1%; Score 857; DB 3; Length 224;

Best Local Similarity 72.9%; Pred. No. 9, 7e-31; Indels 4; Gaps 3;

Matches 164; Conservative 27; Mismatches 30; Indels 4; Gaps 3;

QY 1 MAGEK - GLVLLDFWVSPFGQRVRLAEGKGLPVEYAEEDLMAGSKDRLLRANPVHKIP 58
 1 ||| 1 ||| 1 ||| 1 ||| 1 ||| 1 ||| 1 ||| 1 ||| 1 : 1 ||| 1 ||| 1 ||| 1 ||| 1 ||| 1 |||
 1 MAEKKQGLQQLDFWVSPFGQRCLRLDEKGLAYELEQDL - RNSSELLRANPVHKIP 59
 1 ||| 1 ||| 1 ||| 1 ||| 1 ||| 1 ||| 1 ||| 1 ||| 1 : 1 ||| 1 ||| 1 ||| 1 ||| 1 ||| 1 |||
 59 VLLHDGRAVNESLIILQYLEAEPD - APALLPSDPTYARAQARFWADYVDKKYDCCGSLRW 117
 1 ||| 1 ||| 1 ||| 1 ||| 1 ||| 1 ||| 1 ||| 1 ||| 1 : 1 ||| 1 ||| 1 ||| 1 ||| 1 ||| 1 |||
 Db 60 VLLHDGRPVCLSLVQYLDEAEPAPALLPADPPTYARAQARFWADYVDKLYDCGTRLW 119
 118 KLKGEPQAQAAEMLDILKTDGALDKPFGGDKGTFVDAAFAPPTAWPHSYERGEFS 177
 1 ||| 1 ||| 1 ||| 1 ||| 1 ||| 1 ||| 1 ||| 1 : 1 ||| 1 ||| 1 ||| 1 ||| 1 ||| 1 ||| 1 |||
 Db 120 KLKGDSQAQAAEMVYLRLTLEGALGDPFGGDKGTFVDAALVPTSWFLAYDRGGVS 179
 1 ||| 1 ||| 1 ||| 1 ||| 1 ||| 1 ||| 1 ||| 1 : 1 ||| 1 ||| 1 ||| 1 ||| 1 ||| 1 ||| 1 |||

QY 178 LPEVAKIAAWAKRCGEREVSAKSLSPDKYDFICLLKKYGLIE 222
 1 :
 Db 180 VEKECPRLAAWAKRCERPSVAKNLYPPEKTYDFVGMKRLGIE 224

RESULT 3

US-08-924-759-12

; Sequence 12, Application US/08924759

; GENERAL INFORMATION:

; APPLICANT: MCCONIGLE, BRIAN

; APPLICANT: O'KEEFE, DANIEL

; TITLE OF INVENTION: PLANT GLUTATHIONE-S-TRANSFERASE ENZYMES

; FILE REFERENCE: CL-1128-A

; CURRENT APPLICATION NUMBER: US/09/248,335

; CURRENT FILING DATE: 1999-02-10

; EARLIER APPLICATION NUMBER: 08/924,759

; EARLIER FILING DATE: 1997-September-05

; NUMBER OF SEQ ID NOS: 74

; SOFTWARE: Microsoft Word Version 7.0A

; SEQ ID NO: 12

; LENGTH: 225

; TYPE: PRT

; ORGANISM: maize

US-09-248-335-12

; Sequence 12, Application US/09248335

; GENERAL INFORMATION:

; APPLICANT: MCCONIGLE, BRIAN

; APPLICANT: O'KEEFE, DANIEL

; TITLE OF INVENTION: PLANT GLUTATHIONE-S-TRANSFERASE

; NUMBER OF SEQUENCES: 24

; CORRESPONDENCE ADDRESS:

; ADDRESSEE: E. I. DU PONT DE NEMOURS AND COMPANY

; STREET: 1007 MARKET STREET

; CITY: WILMINGTON

Query Match 69.8%; Score 818.5; DB 3; Length 225;
 Best Local Similarity 63.5%; Pred. No. 2.6e-86;
 Matches 157; Conservative 31; Mismatches 33; Indels 5; Gaps 4;

1 MAGE-KGIVLDFWVSPFGQRVRIALAEKGKLPYEAEEDLMAK-KSDRLLRANPVHKKI 57
 1 MAGETKGVLVLDWVSPFGQRVRIALAEKGKLPYEAEEDLMAK-KSDRLLRANPVHKKI 60

58 PVLLHDGRAYNESTLILQYLEEFAPDA-PALPSDPYARAQAREWADYDKTXYDCGSPU 116
 61 PVLLHDGRAYNESTLILQYLEEFAPDA-PALPSDPYARAQAREWADYDKTXYDCGSPU 119

117 WKLGEPQQARQARAVIVQYRNLLGELGKKAFFGSEAFGSDVDAVLPYFPWLPSYERQDF 176
 120 WKLGEPQQARQARAVIVQYRNLLGELGKKAFFGSEAFGSDVDAVLPYFPWLPSYERQDF 179

RESULT 5
 US-08-924-747-6
 Sequence 6, Application US/08924747
 Patent No. 6063570

GENERAL INFORMATION:
 APPLICANT: MCGONGLE, BRIAN
 APPLICANT: O'KEEFE, DANIEL
 TITLE OF INVENTION: SOYBEAN GLUTATHIONE-S-TRANSFERASE
 TITLE OF INVENTION: ENYMES
 NUMBER OF SEQUENCES: 32

CORRESPONDENCE ADDRESS:
 ADDRESSEE: E.I. DU PONT DE NEMOURS AND COMPANY
 STREET: 1007 MARKET STREET
 CITY: WILMINGTON
 STATE: DELAWARE
 COUNTY: UNITED STATES OF AMERICA
 ZIP: 19898

COMPUTER READABLE FORM:
 MEDIUM TYPE: DISKETTE, 3.50 INCH
 COMPUTER: IBM PC COMPATIBLE
 OPERATING SYSTEM: MICROSOFT WORD FOR WINDOWS 95
 SOFTWARE: MICROSOFT WORD VERSION 7.0A
 CURRENT APPLICATION DATA:
 APPLICATION NUMBER: US/08/924-747

FILING DATE:
 CLASSIFICATION: 435
 ATTORNEY/AGENT INFORMATION:
 NAME: FLOYD, LINDA AXAMETHY
 REGISTRATION NUMBER: 33,692
 REFERENCE/DOCKET NUMBER: CL-1108

TELECOMMUNICATION INFORMATION:
 TELEPHONE: 302-892-8112
 TELEX/FAX: 302-773-0164
 INFORMATION FOR SEQ ID NO: 6:
 SEQUENCE CHARACTERISTICS:
 LENGTH: 219 amino acids

TYPE: amino acid
 STRANDEDNESS: not relevant
 TOPOLOGY: not relevant
 MOLECULE TYPE: protein
 ORIGINAL SOURCE: SOYBEAN
 TISSUE TYPE: SOYBEAN
 IMMEDIATE SOURCE:
 CLONE: GSTA

US-08-924-747-6

Query Match 58.3%; Score 683.5; DB 4; Length 219;
 Best Local Similarity 60.2%; Pred. No. 8.9e-71;
 Matches 130; Conservative 27; Mismatches 58; Indels 1; Gaps 1;

Query Match 58.3%; Score 683.5; DB 4; Length 219;
 Best Local Similarity 60.2%; Pred. No. 8.9e-71;
 Matches 130; Conservative 27; Mismatches 58; Indels 1; Gaps 1;

Query Match 69.8%; Score 818.5; DB 3; Length 225;
 Best Local Similarity 63.5%; Pred. No. 2.6e-86;
 Matches 157; Conservative 31; Mismatches 33; Indels 5; Gaps 4;

5 VLLDDEWVSPFGQRVRIALAEKGKLPYEAEEDLMAK-KSDRLLRANPVHKKI 63
 67 VNESSLILQYLEEFAPDA-PALPSDPYARAQAREWADYDKTXYDCGSPU 126
 64 ICESLIAQYIEWYNDRNPPLPSDPYQRAQTREWADYDKTXYDCGSPU 123
 127 ARAEMLDILKTLGALGDKPFQGGDGFGEVDAAFAPFTAWHESYRGFSLEPEVAKIA 186
 124 ARKEFIEAKLQLEOLQDGTYFGGDNLGVDIAVNPFTWKFAYETFGTLLNIEECPRF 183
 187 AWAKRCGEREVSAKSLSPDKYDFIGLKKYCGIE 222
 184 AWAKRCLOQESVAKSLPQDQKVYEFIMDLRKKLIGIE 219

RESULT 6
 US-09-296-715-6
 Sequence 6, Application US/09296715
 Patent No. 6171839

GENERAL INFORMATION:
 APPLICANT: MCGONGLE, BRIAN
 APPLICANT: O'KEEFE, DANIEL
 TITLE OF INVENTION: SOYBEAN GLUTATHIONE-S-TRANSFERASE
 NUMBER OF SEQUENCES: 32

CORRESPONDENCE ADDRESS:
 ADDRESSEE: E.I. DU PONT DE NEMOURS AND COMPANY
 STREET: 1007 MARKET STREET
 CITY: WILMINGTON
 STATE: DELAWARE
 COUNTY: UNITED STATES OF AMERICA
 ZIP: 19898

COMPUTER READABLE FORM:
 MEDIUM TYPE: DISKETTE, 3.50 INCH
 COMPUTER: IBM PC COMPATIBLE
 OPERATING SYSTEM: MICROSOFT WORD FOR WINDOWS 95
 SOFTWARE: MICROSOFT WORD VERSION 7.0A
 CURRENT APPLICATION DATA:
 APPLICATION NUMBER: US/09/296,715

FILING DATE:
 CLASSIFICATION:
 ATTORNEY/AGENT INFORMATION:
 NAME: FLOYD, LINDA AXAMETHY
 REGISTRATION NUMBER: 33,692
 REFERENCE/DOCKET NUMBER: CL-1108

TELECOMMUNICATION INFORMATION:
 TELEPHONE: 302-892-8112
 TELEX/FAX: 302-773-0164
 INFORMATION FOR SEQ ID NO: 6:
 SEQUENCE CHARACTERISTICS:
 LENGTH: 219 amino acids

TYPE: amino acid
 STRANDEDNESS: not relevant
 TOPOLOGY: not relevant
 MOLECULE TYPE: protein
 ORIGINAL SOURCE: SOYBEAN
 TISSUE TYPE: SOYBEAN
 IMMEDIATE SOURCE:
 CLONE: GSTA

US-09-296-715-6

Query Match 58.3%; Score 683.5; DB 4; Length 219;
 Best Local Similarity 60.2%; Pred. No. 8.9e-71;
 Matches 130; Conservative 27; Mismatches 58; Indels 1; Gaps 1;

7 LYLLDDEWVSPFGQRVRIALAEKGKLPYEAEEDLMAK-KSDRLLRANPVHKKI 66
 5 VLLDDEWVSPFGQRVRIALAEKGKLPYEAEEDLMAK-KSDRLLRANPVHKKI 63
 67 VNESSLILQYLEEFAPDA-PALPSDPYARAQAREWADYDKTXYDCGSPU 126
 64 ICESLIAQYIEWYNDRNPPLPSDPYQRAQTREWADYDKTXYDCGSPU 123

RESULT 12
 US-09-247-373B-12 ; Sequence 12, Application US/0924737B
 Patent No. 616854
 GENERAL INFORMATION:
 APPLICANT: MC GONGIGLE, BRIAN
 TITLE OF INVENTION: SOYBEAN GLUTATHIONE-S-TRANSFERASE ENZYME
 FILE REFERENCE: CL-1108-A
 CURRENT APPLICATION NUMBER: US/09/247, 373B
 CURRENT FILING DATE: 1999-02-10
 PRIOR APPLICATION NUMBER: 08/924, 747
 PRIOR FILING DATE: 1997-09-05
 NUMBER OF SEQ ID NOS: 56
 SOFTWARE: Microsoft Office 97
 SEQ ID NO: 12
 LENGTH: 216
 TYPE: PRT
 ORGANISM: SOYBEAN
 US-09-247-373B-12

Query Match 54.5% ; Score 639, DB 4; Length 2
 Best Local Similarity 55.9% ; Pred. No. 1.2e-59;
 Matches 124; Conservative 33; Mismatches 59; Indels 1

Qy 1 MAGEKGLVILDFWNSPFGQRVRIALAEKGPLYEABDLMAGKSDR
 Db 1 MADE--VYILDFWNSPFGQRVRIALAEKGIKYESKEDLQ_NKSPLI

Qy 61 LHDPGRAVNLSLITLQYLEAFPDAAPLLPSDPYDARAQAFREWADYVDR
 Db 58 IHNKPICSLVAVQYIEEYWNDRNPLPSDPYQRAQAFREWADFDVN

Qy 121 GEPOQAQRGEAEMLDLTKLTDGALGDPKPFGGDKFGFVDAAFAPFTANR
 Db 118 GEEPKAQRKFIEALKLLEFOLQDGLTFGGDDLGPFVDAALIPFDTWR

Qy 181 VAPKIAAKAQRKGGEREVSAKSLSPDKVYDEIGLLKKKYGIE 222
 Db 175 ECEKFVAAKRCRGEREVSAKSLSPDKVYDEIGLLKKKYGIE 216

RESULT 13
 US-09-296-715-12 ; Sequence 12, Application US/09296715
 Patent No. 6171839
 GENERAL INFORMATION:
 APPLICANT: MC GONGIGLE, BRIAN
 APPLICANT: O'KEEFE, DANIEL
 TITLE OF INVENTION: SOYBEAN GLUTATHIONE-S-TRANSFERASE
 TITLE OF INVENTION: ENZYME
 NUMBER OF SEQUENCES: 32
 CORRESPONDENCE ADDRESS:
 ADDRESSEE: E. I. DU PONT DE NEMOURS AND COMPANY
 STREET: 1007 MARKET STREET
 CITY: WILMINGTON
 STATE: DELAWARE
 COUNTRY: UNITED STATES OF AMERICA
 ZIP: 19898
 COMPUTER READABLE FORM:
 MEDIUM TYPE: DISKETTE, 3.50 INCH
 COMPUTER: IBM PC COMPATIBLE
 OPERATING SYSTEM: MICROSOFT WORD FOR WINDOWS 95
 SOFTWARE: MICROSOFT WORD VERSION 7.0A
 CURRENT APPLICATION DATA:
 APPLICATION NUMBER: US/09/296,715
 FILING DATE:
 CLASSIFICATION:
 ATTORNEY/AGENT INFORMATION:
 NAME: FLOYD, LINDA AXAMETHY

SEQUENCE CHARACTERISTICS:
 LENGTH: 200 amino acids
 TYPE: amino acid
 STRANDEDNESS: not relevant
 TOPOLOGY: not relevant
 MOLECULE TYPE: protein
 ORIGINAL SOURCE:
 TISSUE TYPE: SOYBEAN
 IMMEDIATE SOURCE:
 CLONE: SSL.PK0002.F7
 US-08-924-747-18

Query Match 51.3% Score 601.5; DB 3; Length 200;
 Best Local Similarity 55.3%; Pred. No. 2.2e-61;
 Matches 109; Conservative 32; Mismatches 55; Indels 1; Gaps 1;

Qy 6 GIVVLDFTWSPFGVRVIALAEGKSPYAAEDELMAGKSDRLLRANPYHKKIPVLLHDGR 65
 Db 4 GVVLLDTWASMGHRVIALAEGVVEYKEENL-RNKSPLLQMNPIHKKIPVLLHNK 62

Qy 66 AVNESLILQYLERAFDAPALLSDPYARAQAFWADYVDKVYDGSRLWKLKGEPQA 125
 Db 63 PICESAIIYQYIDEWVNDKAPILPSDPIYERAQAFRWWDYIDKVVDTWKMWSKGEEH 122

Qy 126 QARAEMLDILKTLGALGDKPFFGDKFGFVDAFAFPFTAWFHSYERYGEFSLPEVAPKI 185
 Db 123 AGKKEFISIFKQLEETLSDKAYGSDDTGFGLDIGLIPFYSWFTETIGNFKMEECPKL 182

Qy 186 AAWAKRCGGERESTVAKSL 202
 Db 183 VAWAKRCMREAVSKSL 199

RESULT 15

US-09-247-373B-18
 ; Sequence 18, Application US/09247373B
 ; Patent No. 616894
 ; GENERAL INFORMATION:
 ; APPLICANT: MC GONIGLE, BRIAN
 ; APPLICANT: O'KEEFE, DANIEL
 ; TITLE OF INVENTION: SOYBEAN GLUTATHIONE-S-TRANSFERASE ENZYMES
 ; FILE REFERENCE: CL-1108-A
 ; CURRENT APPLICATION NUMBER: US/09/247,373B
 ; CURRENT FILING DATE: 1999-02-10
 ; PRIOR APPLICATION NUMBER: 08/924,747
 ; PRIOR FILING DATE: 1997-09-05
 ; NUMBER OF SEQ ID NOS: 56
 ; SOFTWARE: Microsoft Office 97
 ; SEQ ID NO 18
 ; LENGTH: 200
 ; TYPE: PRT
 ; ORGANISM: SOYBEAN
 US-09-247-373B-18

Query Match 51.3% Score 601.5; DB 4; Length 200;
 Best Local Similarity 55.3%; Pred. No. 2.2e-61;
 Matches 109; Conservative 32; Mismatches 55; Indels 1; Gaps 1;

Qy 6 GIVVLDFTWSPFGVRVIALAEGKSPYAAEDELMAGKSDRLLRANPYHKKIPVLLHDGR 65
 Db 4 GVVLLDTWASMGHRVIALAEGVVEYKEENL-RNKSPLLQMNPIHKKIPVLLHNK 62

Qy 66 AVNESLILQYLERAFDAPALLSDPYARAQAFWADYVDKVYDGSRLWKLKGEPQA 125
 Db 63 PICESAIIYQYIDEWVNDKAPILPSDPIYERAQAFRWWDYIDKVVDTWKMWSKGEEH 122

Qy 126 QARAEMLDILKTLGALGDKPFFGDKFGFVDAFAFPFTAWFHSYERYGEFSLPEVAPKI 185
 Db 123 AGKKEFISIFKQLEETLSDKAYGSDDTGFGLDIGLIPFYSWFTETIGNFKMEECPKL 182

Qy 186 AAWAKRCGGERESTVAKSL 202
 Db 183 VAWAKRCMREAVSKSL 199

GenCore version 5.1.6
 Copyright (c) 1993 - 2003 Compugen Ltd.

DB protein - protein search, using sw model
 Run on: July 1, 2003, 08:53:35 ; Search time 21 Seconds
 (without alignments)
 1159.124 Million cell updates/sec

Protein score: 1173
 Perfect score: 1173
 Sequence: 1 MAGERGLVLLDFWWSPFQQR.....YSPDKVYDFGLLKKYKIE 222

Scoring table: BLOSUM62
 Gapext 10.0 , Gapext 0.5
 424699 seqs, 103646833 residues

Searched: 424699 seqs, 103646833 residues
 Total number of hits satisfying chosen parameters: 424699

Minimum DB seq length: 0
 Maximum DB seq length: 2000000000
 Post-processing: Minimum Match 0%

ALIGNMENTS

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

EXAMPLES

Result No.	Score	Query Match	Length	DB ID	Description	
					%	Sequence
1	405.5	34.6	236	10	US-09-765-213A-9	Sequence 9, Appli
2	362.5	30.9	229	10	US-09-765-213A-8	Sequence 8, Appli
3	334	28.5	233	10	US-09-765-213A-7	Sequence 2, Appli
4	331.5	28.3	225	10	US-09-765-213A-7	Sequence 7, Appli
5	320.5	27.3	224	10	US-09-765-213A-11	Sequence 11, Appli
6	319	27.2	233	10	US-09-765-213A-4	Sequence 4, Appli
7	309	26.3	233	10	US-09-765-213A-6	Sequence 6, Appli
8	291	24.8	240	10	US-09-765-213A-10	Sequence 10, Appli
9	235	20.0	83	10	US-09-050-010-7	Sequence 7, Appli
10	198	16.9	212	10	US-09-741-669-351	Sequence 351, App
11	172.5	14.7	241	9	US-09-854-133-194	Sequence 134, App
12	172.5	14.7	241	9	US-10-225-273-4	Sequence 4, Appli
13	172.5	14.7	241	10	US-09-738-973-194	Sequence 194, App
14	172.5	14.7	286	10	US-09-935-302-653	Sequence 653, App
15	167.5	14.3	241	9	US-09-854-133-204	Sequence 204, App
16	167.5	14.3	241	10	US-09-738-973-204	Sequence 204, App
17	129.5	11.0	86	10	US-09-050-010-6	Sequence 6, Appli
18	124	10.6	195	10	US-09-798-164	Sequence 164, App
19	121.5	10.4	263	9	US-09-769-787-168	Sequence 68, Appli

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

RESULT 2
US-09-765-213A-8
; Sequence 8, Application US/09765213A
; Patent No. US20020079846A1
; GENERAL INFORMATION:
; APPLICANT: Faccioli, Peter J
; TITLE OF INVENTION: No. US20020079846A1 Glutathione-S-Transferase Nucleic Acids and Methods of Use
; TITLE OF INVENTION: Polypeptides and Methods of Use
; FILE REFERENCE: 22542-001
; CURRENT FILING DATE:
; CURRENT APPLICATION NUMBER: US/09/765,213A
; PRIOR FILING DATE: 2001-01-17
; NUMBER OF SEQ ID NOS: 11
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 8
; LENGTH: 229
; TYPE: PRT
; ORGANISM: Aegilops squarrosa
US-09-765-213A-8

Query Match 30.9%; Score 362.5; DB 10; Length 229;
Best Local Similarity 42.3%; Pred. No. 2.5e-10;
Matches 93; Conservative 33; Mismatches 79; Indels 15; Gaps 8;

Qy 2 AGBKGLVLLDFTWVSPGQVRVIALAERKGLPKEYAEDDLMAKSDRLLRANPVHKKIPV 61
Db 3 AGGDLKLLGAWPSPEVTRKIALALKLSSVEDVEEDLYK KSELLIKSNPEVHKKIPV 61
Qy 62 HDGRAYNESLILQLYEEAFAFPD-APALLPSPDYARAQARFADYYDKKVVQGSRLWKLK 120
Db 62 HNGAPICESMILQLYIDDEVFAGTGPSLIPADYERARFWAVDDKLV--APWROW-LR 118

Qy 121 G---EPOQAQARAEMLDILKTDLGALGD---KPFEGGDKEFVDAFAPFTAWFHSYERY 173
Db 119 GRTEEERSEGGKQAFAAVGVLLEGALRECCKSGCGFFGCGDVG-LDVALGGVLSWMKYTEAL 177

Qy 174 GESLPLVVA--PKIAAWAKRCGERESVAKSYSPDVKYDF 211
Db 178 SGDKIFDAAKTPLLAVMVERFTELDRAKAALPDVGRLLEF 217

RESULT 3
US-09-765-213A-2
; Sequence 2, Application US/09765213A
; Patent No. US20020079846A1
; GENERAL INFORMATION:
; APPLICANT: Faccioli, Peter J
; TITLE OF INVENTION: No. US20020079846A1 Glutathione-S-Transferase Nucleic Acids and Methods of Use
; TITLE OF INVENTION: Polypeptides and Methods of Use
; FILE REFERENCE: 22542-001
; CURRENT FILING DATE:
; CURRENT APPLICATION NUMBER: US/09/765,213A
; PRIOR FILING DATE: 2001-01-17
; NUMBER OF SEQ ID NOS: 11
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 2
; LENGTH: 233
; TYPE: PRT
; ORGANISM: Papaver somniferum
US-09-765-213A-2

Query Match 28.5%; Score 334; DB 10; Length 233;
Best Local Similarity 35.1%; Pred. No. 2.7e-27;
Matches 81; Conservative 42; Mismatches 88; Indels 20; Gaps 7;

Qy 2 AGEKGLVLLDFTWVSPGQVRVIALAERKGLPKEYAEDDLMAKSDRLLRANPVHKKIPV 61
Db 4 SSSEEEVKGGMPSPEVWMPRTALNPKSVKYLLEETFGSKSELLIKSNPEVHKKIPV 62
Qy 62 HDGRAYNESLILQLYEEAFAFPD-APALLPSPDYARAQARFADYYDKKVVQGSRLWKLK 120

RESULT 4
US-09-765-213A-7
; Sequence 7, Application US/09765213A
; Patent No. US20020079846A1
; GENERAL INFORMATION:
; APPLICANT: Faccioli, Peter J
; TITLE OF INVENTION: No. US20020079846A1 Glutathione-S-Transferase Nucleic Acids and Methods of Use
; FILE REFERENCE: 22542-001
; CURRENT APPLICATION NUMBER: US/09/765,213A
; CURRENT FILING DATE: 2001-01-17
; PRIOR APPLICATION NUMBER: 60/176708
; PRIOR FILING DATE: 2000-01-18
; NUMBER OF SEQ ID NOS: 11
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 7
; LENGTH: 225
; TYPE: PRT
; ORGANISM: Arabidopsis thaliana
US-09-765-213A-7

Query Match 28.3%; Score 331.5; DB 10; Length 225;
Best Local Similarity 34.7%; Pred. No. 4.7e-17;
Matches 82; Conservative 46; Mismatches 83; Indels 25; Gaps 9;

Qy 1 MAGEKGLVLLDFTWVSPGQVRVIALAERKGLPKEYAEDDLMAKSDRLLRANPVHKKIPV 59
Db 1 MRQNDTYKLIGWSSSPSLRARVALHKSVKYELDDEPVKEKSSELLIKSNPIHKKIPV 60
Qy 60 LLHDGRAYNESLILQLYEEAFAFPDAPALLPSPDYARAQARFADYYDKKVVQGSRLWKLK 119
Db 61 LLHGDLISSELSLNVQQ-DEAWPSVSPSILPSDAYRASAREFQIDDK---CFAAVDAV 115

Qy 120 KG---EPOQAQARAEMLDILKTDLTD---GALGDKPFEGGDKEFVDAFAPFTAWFHF 168
Db 116 VGAKDDGKMAVGKLMCILATEETQKSSRGGLG--FEGETICITYLDRCSALGPPI 172

Qy 169 SYERYG--EFSLPEVARKIAAWAKRCGERESVAKSYSPDVKYDFGLLKKYKIE 222
Db 173 VTEAFSCVKFLQETTFLGLIWAERFRAHEAVKPF--YMP-TVEEVYAFAKOKEVQ 225

RESULT 5
US-09-765-213A-11
; Sequence 11, Application US/09765213A
; Patent No. US20020079846A1
; GENERAL INFORMATION:
; APPLICANT: Faccioli, Peter J
; TITLE OF INVENTION: No. US20020079846A1 Glutathione-S-Transferase Nucleic Acids and Methods of Use
; FILE REFERENCE: 22542-001
; CURRENT APPLICATION NUMBER: US/09/765,213A
; CURRENT FILING DATE: 2001-01-17
; PRIOR APPLICATION NUMBER: 60/176708
; PRIOR FILING DATE: 2000-01-18
; NUMBER OF SEQ ID NOS: 11
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 11
; LENGTH: 224
; TYPE: PRT
; ORGANISM: Glycine max
US-09-765-213A-11

Query Match 27.3%; Score 320.5; DB 10; Length 224;
 Best Local Similarity 35.1%; Pred. No. 6.8e-26;
 Matches 78; Conservative 49; Mismatches 84; Indels 11; Gaps 7;
 SEQ ID NO 6

QY 2 AGEKGLVNLDEWWNSPFGQVRVIALAEKGPLYEAAEDLMAGKSDRLLRANPVHKKIPYL 61
 Db 3 ATQDVKLGLGIVGQFVQIAKLKGPEYKFLIEENL-GNKSDRLLKLYNPVKKVKPVY 61
 LENGTH: 233

QY 62 HDGRAVNESLTILQYLEEAFPDAALLSPDPYARAQAFWADYVDKKPYDCGSR-LWK 120

Db 62 HNEQPIAESLIVVYIDEWKNNP-ILPSPDQYQALARWSKFKDDKIVGAVSKSVFTID 120

QY 121 GEPAQARAEMLDILKTLGDKPFEGDKFGF-VDAAFAFPFTAWFHSYERYGEFSI- 178

Db 121 EKEREKNEVETYEALQFLNELDKDKFGEFFGLDIAVFTIAH--WIFIFOETAGLQLF 178

QY 179 -PEVAPKIAAWAKRCGERRSVAKSLYSPDKVYDFIGLKKY 219

Db 179 TSEKEPILKWSQEFPLNHPVHEVLPPLPDLFAY--FRLARY 217

RESULT 6

US 09-765-213A-4

Sequence 4, Application US/09765213A
 Patent No. US20020073846A1

GENERAL INFORMATION:

APPLICANT: Faccini, Peter J

TITLE OF INVENTION: No. US20020079846A1 *Glutathione-S-Transferase Nucleic Acids and Methods of Use*

FILE INVENTION: Polypeptides and Methods of Use

CURRENT APPLICATION NUMBER: US/09/765,213A

CURRENT FILING DATE: 2001-01-17

PRIOR APPLICATION NUMBER: 60/176708

PRIOR FILING DATE: 2000-01-18

SEQ ID NO: 4

SOFTWARE: PatentIn Ver. 2.1

LENGTH: 233

TYPE: PRF

ORGANISM: Papaver somniferum

US 09-765-213A-4

Query Match 27.2%; Score 319; DB 10; Length 233;
 Best Local Similarity 33.8%; Pred. No. 1e-23; Mismatches 90; Indels 20; Gaps 7;
 Matches 78; Conservative 43; Mismatches 90; Indels 20; Gaps 7;

QY 2 AGEKGLVNLDEWWNSPFGQVRVIALAEKGPLYEAAEDLMAGKSDRLLRANPVHKKIPYL 61
 Db 4 SGSEEVKILGGWPSPFVMPRILNIKSKY-YLLEETFGSKSLLKSNPIKKIPYL 62

QY 62 HDGRAVNESLTILQYLEEAFPDA-PALLSPDPYARAQAFWADYVDKKPYDCGSR-LWK 120
 Db 63 HDKDPICEEMIIVYDWDWASAGHSIISDPYDASIAFWRAYIDDKFPLSMLGIASK 122

QY 121 GEPAQARAEMLDILKTLGDKPFEGDKFGF-VDAAFAFPFTAWFHSYI- 171
 Db 123 DAEERKKAEEQATIAAFGILLEEAYQTSKKG---KTLDALGDKPFEGDKFGF-VDAAFAFPFTAWFHSYI- 171

QY 172 RYGEFSL--PEVAPKIAAWAKRCGERRSVAKSLYSPDKVYDFIGLKKY 220
 Db 179 KMNGIKLFDEXXKVFGLTWAEKFCADETVKSVMPETDALMEF--AKKFG 226

RESULT 8

US 09-765-213A-10

Sequence 10, Application US/09765213A
 Patent No. US20020073846A1

GENERAL INFORMATION:

APPLICANT: Faccini, Peter J

TITLE OF INVENTION: No. US20020079846A1 *Glutathione-S-Transferase Nucleic Acids and Methods of Use*

FILE REFERENCE: 22542-001

CURRENT APPLICATION NUMBER: US/09/765,213A

PRIOR APPLICATION NUMBER: 60/176708

PRIOR FILING DATE: 2000-01-18

NUMBER OF SEQ ID NOS: 11
; SOFTWARE: Patentin Ver. 2.1
; SEQ ID NO 10
; LENGTH: 240
; TYPE: PRT
; ORGANISM: Zea mays
; US-09-765-213A-10

Query Match 24 8%; Score 291, DB 10; Length 240;
Best Local Similarity 36.8%; Pred. No. 9.9e-23; Mismatches 81; Conservative 31; Indels 18; Gaps 7;

QY 14 VSPFGQRVRAALAEGKLPYIAYEEDLMAGSKDRLLRANPVHKKIPV-LIHGRAYNESLT 72
Db 13 VSPFGQRVRAALAEGKLPYIAYEEDLMAGSKDRLLRANPVHKKIPV-PVLLIDGRACESAV 72

QY 73 ILQYLEAFTPDA-----PALLPSDPAQRARQAFWADYYDKVYDGSRILWKLKGEP--Q 124
Db 73 IVOQYIEDVARESGGAEAGSLLPDPDYERAMHRWTFATIDKFPALDAY-SLAFTPGAR 131

QY 125 AQAAREMFLDLIKTUDGALGSKP----FFGDKFGEFYDAFAAFAFTAWHSHYERYGEFL 178
Db 132 AQAATRALSLLBEAKFORSNGRAFFSGDAAPGLLDAALGCFLPALRACERIHLGLS 191

QY 179 PEVA--PKIAAWAKRCGERRSVAKSLSYSPKVDYFIGLK 216
Db 192 IDASATPLLGWSQRFAAHAAKRVLPDTKVVQFTRLQ 231

RESULT 9
US-09-050-010-7
; Sequence 7, Application US/09050010
; Patent No. US2001001003A1
; GENERAL INFORMATION:
; APPLICANT: Nagai, Keiichi
; APPLICANT: Irie, Ryotaro
; APPLICANT: Hiraoka, Susumu
; APPLICANT: Kasahara, Naoko
; TITLE OF INVENTION: METHOD FOR COMPARISON OF DNA BASE
; NUMBER OF SEQUENCES: 12
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: FAY, SHARPE, BEALL, FAGAN, MINNICH & MCKEE
; STREET: 104 East Hume Avenue
; CITY: Alexandria
; STATE: Virginia
; COUNTRY: USA
; ZIP: 22301
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.30
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US09/0500,010
; FILING DATE: 30-MAR-1998
; CLASSIFICATION: 435
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: JP 09-079586
; ATTORNEY/AGENT INFORMATION:
; NAME: Beall, Jr., Thomas E.
; REGISTRATION NUMBER: 22,410
; REFERENCE/DOCKET NUMBER: ASA-707
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 703-384-1120
; TELEFAX: 703-884-1167
; INFORMATION FOR SEQ ID NO: 7:
; LENGTH: 83 amino acids
; TYPE: amino acid
; STRANDEDNESS:
; TOPOLOGY: linear

; MOLECULE TYPE: protein
; US-09-050-010-7
; Query Match 20 0%; Score 235; DB 10; Length 83;
; Best Local Similarity 59.5%; Pred. No. 1.9e-17; Mismatches 13; Indels 2; Gaps 2;
; Matches 50; Conservative 19; Indels 2; Gaps 2;

QY 1 MAGE KGLIVLDEWTSPEGVRIALAAEKGLPYIAYEEDLMAGSKDRLLRANPVHKKIPV 59
Db 1 MAGSGDELMILGKWNSPFTRVELALGKLSYEVKQDLY-NKSELLLASNPVHKKIPV 59

QY 60 LLHDGRAVNESLTLIQLQYLEAFTPDI 83
Db 60 LIHNEKPVCESSIVYIYBAPFD 83

RESULT 10
US-09-741-669-351
; Sequence 351, Application US/09741669
; Patent No. US20030022718A1
; GENERAL INFORMATION:
; APPLICANT: Forsyth, R. Allyn
; APPLICANT: Ohleen, Kari L.
; APPLICANT: Elyskind, Judith W.
; APPLICANT: Title of Invention: Genes identified as required for
; proliferation of E. coli
; FILE REFERENCE: ELTRA.009A
; CURRENT APPLICATION NUMBER: US/09/741,669
; CURRENT FILING DATE: 2000-12-19
; PRIOR APPLICATION NUMBER: US 60/173005
; NUMBER OF SEQ ID NOS: 481
; SEQ ID NO: 351
; SOFTWARE: FastSEQ for Windows Version 4.0
; LENGTH: 212
; TYPE: PRT
; ORGANISM: Escherichia coli
; US-09-741-669-351

Query Match 16 9%; Score 198; DB 10; Length 212;
Best Local Similarity 30.9%; Pred. No. 5.8e-13; Mismatches 36; Indels 12; Gaps 8;
Matches 60; Conservative 36; Indels 12; Gaps 8;

QY 17 FGQYRVTIAEKGIPY--EYAEEDLMAGSKDRLLRANPVHKKIPV-LIHGRAYNESLIL 74
Db 21 YSHQVRIVLAEGKYSFEIEHVEKD--NPFQDLIDLN-P-NQSVPVTLVDRLETLWESRILM 76

QY 75 QYLEFAFPDAPALLPSDPYKARAQAFRWDAYDKVYDCCSRILWKLKGEPQAQARAEMLDI 134
Db 77 EYLDEREPH-PLMMPYVARGESRMLHRIEKWYTLMTNTIINGASERDARKQLEEE 135

QY 135 LKTLDGALGDKPFFGDKFGEVDAFAFPFTAWHSHYERYG-EFSLPEVAPKIAAWAKRCG 193
Db 136 LLAAAPVFGOKPYFSLDEFSLVDCYLAPL-LW-RLPQLGIEFSGPG-AKELKGYMTRYF 191

QY 194 ERESYAKSLYSYSPDK 207
Db 192 ERDSFLASTEAER 205

RESULT 11
US-09-854-133-194
; Sequence 194, Application US/09854133
; Publication No. US2002018349A1
; GENERAL INFORMATION:
; APPLICANT: Lodes, Michael J.
; APPLICANT: Mohamath, Raodoh
; APPLICANT: Henderson, Robert A.
; APPLICANT: Benson, Darin R.
; APPLICANT: Sechrist, Heather
; TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR
; TITLE OF INVENTION: THE THERAPY AND DIAGNOSIS OF LUNG CANCER
; FILE REFERENCE: 210121.475C10

CURRENT APPLICATION NUMBER: US/09/854,133
; CURRENT FILING DATE: 2001-05-11
; NUMBER OF SEQ ID NOS: 735
; SOFTWARE: fastSEQ for Windows version 3.0
; SEQ ID NO: 194
; LENGTH: 241
; TYPE: PRT
; ORGANISM: Homo sapien
; SEQ ID NO: 194-133-194

Query Match 14.7%; Score 172.5; DB 9; Length 241;
Best Local Similarity 28.7%; Pred. No. 3.4e-10;
Matches 58; Conservative 28; Mismatches 107; Indels 9; Gaps 6;

QY 16 PFGQRVIALAEGKGLPEAYEEDLMACKSDRLLRANPVHKKIPVLLH-DGRAVNESSLIL 74
Db 33 PPAERTVLKARGIRHEVINVNL-KURKPEFEKKNP-FGLYPVLENQGOLIYESAATC 90

QY 75 QYLEEAPPDAALLPSPDYARQARFADYVDRKVKYDGSRILWKLKGEPQAQRAEMLDI 134
Db 91 EYLDEAYP-GKLLLPDPYEAQCKQMILELFSKVPVLGSFIRSQNKEGYAGIKEEKFKE 149

QY 135 LRTLDGAQDK-PFFGGDKGFVDAAFAPFTAWHESYERYGEFSLPEVAPKIAAWAKRC 192
Db 150 FTKLEEVLTNKKTFFGNSISMDYLWIP--WFERLEAMKLNECVDHTPKLKLWMAAM 206

QY 193 GEREYVAKSLYSPDKYDFIGL 214
Db 207 KEDPTVSALLTSEKDNGFEL 228

RESULT 13
US-09-738-973-194
; Sequence 194, Application US/09/854,133
; Patent No. US20020110563A1
; GENERAL INFORMATION:
; APPLICANT: Reed, Steven G.
; APPLICANT: Henderson, Robert A.
; APPLICANT: Lodes, Michael J.
; APPLICANT: Fling, Steven P.
; APPLICANT: Mohanath, Raodoh
; APPLICANT: Algata, Paul A.
; APPLICANT: Sechrist, Heather
; APPLICANT: Indrietas, Carol Joseph
; APPLICANT: Penson, Darin R.
; APPLICANT: Elliott, Mark
; APPLICANT: Mannion, Jane
; APPLICANT: Klos, Michael D.
; TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR
; TREATMENT OF CANCER
; TITLE OF INVENTION: THE THERAPY AND DIAGNOSIS OF LUNG CANCER
; FILE REFERENCE: 210121.4759
; CURRENT APPLICATION NUMBER: US/09/738,973
; CURRENT FILING DATE: 2000-12-14
; NUMBER OF SEQ ID NOS: 587
; SOFTWARE: FastSEQ for Windows Version 3.0
; SEQ ID NO: 194
; LENGTH: 241
; TYPE: PRT
; ORGANISM: Homo sapien
; SEQ ID NO: 194-133-194

Query Match 14.7%; Score 172.5; DB 10; Length 241;
Best Local Similarity 28.7%; Pred. No. 3.4e-10;
Matches 58; Conservative 28; Mismatches 107; Indels 9; Gaps 6;

QY 16 PFGQRVIALAEGKGLPEAYEEDLMACKSDRLLRANPVHKKIPVLLH-DGRAVNESSLIL 74
Db 33 PFAERTVLKARGIRHEVINVNL-KNKPENFEKKNP-FGLYPVLENQGOLIYESAATC 90

QY 75 QYLEEAPPDAALLPSPDYARQARFADYVDRKVKYDGSRILWKLKGEPQAQRAEMLDI 134
Db 91 EYLDEAYP-GKLLLPDPYEAQCKQMILELFSKVPVLGSFIRSQNKEGYAGIKEEKFKE 149

QY 135 LRTLDGAQDK-PFFGGDKGFVDAAFAPFTAWHESYERYGEFSLPEVAPKIAAWAKRC 192
Db 150 FTKLEEVLTNKKTFFGNSISMDYLWIP--WFERLEAMKLNECVDHTPKLKLWMAAM 206

QY 193 GEREYVAKSLYSPDKYDFIGL 214
Db 207 KEDPTVSALLTSEKDNGFEL 228

RESULT 14
US-09-225-273-4
; Sequence 4, Application US/10225273
; Publication No. US2003002721A1
; GENERAL INFORMATION:
; APPLICANT: Pfizer Inc.
; APPLICANT: Gabel, Christopher A.
; APPLICANT: Dombroski, Mark A.
; APPLICANT: Geoghegan, Kieran
; APPLICANT: Griffiths, Richard J.
; APPLICANT: Eggerle, James F.
; TITLE OF INVENTION: DIARYLSULFONYLUREA BINDING PROTEINS
; FILE REFERENCE: PC98603
; CURRENT APPLICATION NUMBER: US/10/225,273
; CURRENT FILING DATE: 2002-08-21
; PRIOR APPLICATION NUMBER: 09/387,372
; PRIOR FILING DATE: 1999-08-21
; NUMBER OF SEQ ID NOS: 8
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO: 4
; LENGTH: 241
; TYPE: PRT
; ORGANISM: Homo Sapiens
; SEQ ID NO: 194-133-194

Query Match 14.7%; Score 172.5; DB 9; Length 241;
Best Local Similarity 28.7%; Pred. No. 3.4e-10;
Matches 58; Conservative 28; Mismatches 107; Indels 9; Gaps 6;

QY 16 PFGQRVIALAEGKGLPEAYEEDLMACKSDRLLRANPVHKKIPVLLH-DGRAVNESSLIL 74
Db 33 PPAERTVLKARGIRHEVINVNL-KURKPEFEKKNP-FGLYPVLENQGOLIYESAATC 90

QY 75 QYLEEAPPDAALLPSPDYARQARFADYVDRKVKYDGSRILWKLKGEPQAQRAEMLDI 134
Db 91 EYLDEAYP-GKLLLPDPYEAQCKQMILELFSKVPVLGSFIRSQNKEGYAGIKEEKFKE 149

QY 135 LRTLDGAQDK-PFFGGDKGFVDAAFAPFTAWHESYERYGEFSLPEVAPKIAAWAKRC 192
Db 150 FTKLEEVLTNKKTFFGNSISMDYLWIP--WFERLEAMKLNECVDHTPKLKLWMAAM 206

QY 193 GEREYVAKSLYSPDKYDFIGL 214
Db 207 KEDPTVSALLTSEKDNGFEL 228

RESULT 15
US-09-925-302-653
; Sequence 653, Application US/09925302
; GENERAL INFORMATION:
; APPLICANT: Rosen et al.
; TITLE OF INVENTION: Nucleic Acids, Proteins and Antibodies
; FILE REFERENCE: PA104
; CURRENT APPLICATION NUMBER: US/09/925,302
; CURRENT FILING DATE: 2001-08-10
; PRIOR APPLICATION NUMBER: PCT/US00/05518
; PRIOR FILING DATE: 2000-03-08
; PRIOR APPLICATION NUMBER: 60/124,270
; PRIOR FILING DATE: 1999-03-12
; NUMBER OF SEQ ID NOS: 896
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO: 653
; LENGTH: 285

